

## RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.**

Application Serial Number: 10/567,365  
Source: IFWP  
Date Processed by STIC: 2/13/06

# ***ENTERED***



JFWP

## RAW SEQUENCE LISTING

DATE: 02/13/2006

PATENT APPLICATION: US/10/567,365

TIME: 12:42:56

Input Set : A:\CorrectedSEQ\_list P770PC00.ST25.txt

Output Set: N:\CRF4\02132006\J567365.raw

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5 <110> APPLICANT: ENKAM Pharmaceutical A/S
9 <120> TITLE OF INVENTION: Compounds comprising LPA
13 <130> FILE REFERENCE: P 770 PC00
C--> 17 <140> CURRENT APPLICATION NUMBER: US/10/567,365
C--> 17 <141> CURRENT FILING DATE: 2006-02-07
17 <160> NUMBER OF SEQ ID NOS: 146
21 <170> SOFTWARE: PatentIn version 3.1
25 <210> SEQ ID NO: 1
27 <211> LENGTH: 15
29 <212> TYPE: PRT
31 <213> ORGANISM: Artificial sequence
35 <220> FEATURE:
37 <223> OTHER INFORMATION: NCAM Fn III, 2 [Swiss-Prot: P13591]: FGFR binding motif
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41 Glu Val Tyr Val Val Ala Glu Asn Gln Gln Gly Lys Ser Lys Ala
42 1          5          10          15
45 <210> SEQ ID NO: 2
47 <211> LENGTH: 16
49 <212> TYPE: PRT
51 <213> ORGANISM: Artificial sequence
55 <220> FEATURE:
57 <223> OTHER INFORMATION: Interleukin-6 receptor beta chain [Swiss-Prot: Q00560]: FGFR
bind
58      ing motif
60 <400> SEQUENCE: 2
62 Asn Ile Glu Val Trp Val Glu Ala Glu Asn Ala Leu Gly Lys Lys Val
63 1          5          10          15
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68 <211> LENGTH: 14
70 <212> TYPE: PRT
72 <213> ORGANISM: Artificial sequence
76 <220> FEATURE:
78 <223> OTHER INFORMATION: Heparan sulfate proteoglycan perlecan [Swiss-Prot: P98160]:
FGFR
79      binding motif
81 <400> SEQUENCE: 3
83 Ala Thr Asn Arg Gln Gly Lys Val Lys Ala Phe Ala His Leu
84 1          5          10
87 <210> SEQ ID NO: 4
89 <211> LENGTH: 16
91 <212> TYPE: PRT
93 <213> ORGANISM: Artificial sequence
97 <220> FEATURE:
99 <223> OTHER INFORMATION: Disintegrin and metalloprotease domain 8 (ADAM-8) [Swiss-
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Prot: Q0

100

5910]: FGFR binding motif

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102 <400> SEQUENCE: 4  
 104 Arg Tyr Val Glu Leu Tyr Val Val Ala Asp Ser Gln Glu Phe Gln Lys  
 105 1 5 10 15  
 108 <210> SEQ ID NO: 5  
 110 <211> LENGTH: 13  
 112 <212> TYPE: PRT  
 114 <213> ORGANISM: Artificial sequence  
 118 <220> FEATURE:  
 120 <223> OTHER INFORMATION: Axonal-associated cell adhesion molecule [NCBI: NP\_446331]:

FGFR

121 binding motif  
 123 <400> SEQUENCE: 5  
 125 Val Ala Glu Asn Ser Arg Gly Lys Asn Val Ala Lys Gly  
 126 1 5 10  
 129 <210> SEQ ID NO: 6  
 131 <211> LENGTH: 14  
 133 <212> TYPE: PRT  
 135 <213> ORGANISM: Artificial sequence  
 139 <220> FEATURE:  
 141 <223> OTHER INFORMATION: Myelin-associated glycoprotein (MAG) [Swiss-Prot: P20917]:

FGFR b

142 binding motif  
 144 <400> SEQUENCE: 6  
 146 Gly Glu Tyr Trp Cys Val Ala Glu Asn Gln Tyr Gly Gln Arg  
 147 1 5 10  
 150 <210> SEQ ID NO: 7  
 152 <211> LENGTH: 14  
 154 <212> TYPE: PRT  
 156 <213> ORGANISM: Artificial sequence  
 160 <220> FEATURE:  
 162 <223> OTHER INFORMATION: FIII,1 domain of NCAM [Swiss-Prot: P13591]: FGFR binding

motif

164 <400> SEQUENCE: 7  
 166 Arg Leu Ala Ala Leu Asn Gly Lys Gly Leu Gly Glu Ile Ser  
 167 1 5 10  
 171 <210> SEQ ID NO: 8  
 173 <211> LENGTH: 16  
 175 <212> TYPE: PRT  
 177 <213> ORGANISM: Artificial sequence  
 181 <220> FEATURE:  
 183 <223> OTHER INFORMATION: Neuronal nicotinic acetylcholine receptor alpha 3 subunit

(CHRNA

184 3) [Swiss-Prot: Q8VHH6/P04757:/Q8R4G9/P32297]: FGFR binding motif  
 186 <400> SEQUENCE: 8  
 188 Lys Tyr Ile Ala Glu Asn Met Lys Ala Gln Asn Val Ala Lys Glu Ile  
 189 1 5 10 15  
 192 <210> SEQ ID NO: 9  
 194 <211> LENGTH: 14  
 196 <212> TYPE: PRT  
 198 <213> ORGANISM: Artificial sequence  
 202 <220> FEATURE:  
 204 <223> OTHER INFORMATION: FIII,1 domain of NCAM (Swiss-Prot: P13591): FGFR binding

motif

206 <400> SEQUENCE: 9

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208 Thr Ile Met Gly Leu Lys Pro Glu Thr Arg Tyr Ala Val Arg
209 1          5          10
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214 <211> LENGTH: 14
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218 <213> ORGANISM: Artificial sequence
222 <220> FEATURE:
224 <223> OTHER INFORMATION: Granulocyte colony stimulating factor receptor precursor (G-
CSF-
225      R; CD114 antigen) [Swiss-Prot: Q99062]: FGFR binding motif
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229 Lys Gly Leu Gly Glu Ile Ser Ala Ala Thr Glu Phe Lys Thr
230 1          5          10
233 <210> SEQ ID NO: 11
235 <211> LENGTH: 13
237 <212> TYPE: PRT
239 <213> ORGANISM: Artificial sequence
243 <220> FEATURE:
245 <223> OTHER INFORMATION: NCAM Fn III, 1 [Swiss-Prot: P13591]: FGFR binding motif
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249 Asn Met Gly Ile Trp Val Gln Ala Glu Asn Ala Leu Gly
250 1          5          10
253 <210> SEQ ID NO: 12
255 <211> LENGTH: 10
257 <212> TYPE: PRT
259 <213> ORGANISM: Artificial sequence
263 <220> FEATURE:
265 <223> OTHER INFORMATION: Granulocyte colony stimulating factor receptor precursor (G-
CSF-
266      R; CD114 antigen) [Swiss-Prot: P40223]: FGFR binding motif
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270 Ile Trp Val Gln Ala Glu Asn Met Leu Gly
271 1          5          10
274 <210> SEQ ID NO: 13
276 <211> LENGTH: 11
278 <212> TYPE: PRT
280 <213> ORGANISM: Artificial sequence
284 <220> FEATURE:
286 <223> OTHER INFORMATION: Cytokine-like factor-1 precursor (CLF-1) [Swiss-Prot:
075462]: FG
287      FR binding motif
289 <400> SEQUENCE: 13
291 Glu Ile Trp Val Glu Ala Thr Asn Arg Leu Gly
292 1          5          10
295 <210> SEQ ID NO: 14
297 <211> LENGTH: 10
299 <212> TYPE: PRT
301 <213> ORGANISM: Artificial sequence
305 <220> FEATURE:
307 <223> OTHER INFORMATION: Interleukin-23 receptor (IL-23R) [Q8NFQ9]: FGFR binding
motif
309 <400> SEQUENCE: 14
311 Val Trp Val Gln Ala Ala Asn Ala Leu Gly

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312 1          5          10
315 <210> SEQ ID NO: 15
317 <211> LENGTH: 13
319 <212> TYPE: PRT
321 <213> ORGANISM: Artificial sequence
325 <220> FEATURE:
327 <223> OTHER INFORMATION: Complement factor 1 q , alpha polypeptide (C1QA) [Swiss-
Prot: Q9D
328      CM6]: FGFR binding motif
330 <400> SEQUENCE: 15
332 Glu Val Trp Ile Glu Lys Asp Pro Ala Lys Gly Arg Ile
333 1          5          10
336 <210> SEQ ID NO: 16
338 <211> LENGTH: 14
340 <212> TYPE: PRT
342 <213> ORGANISM: Artificial sequence
346 <220> FEATURE:
348 <223> OTHER INFORMATION: Fasciclin II precursor (FAS2) [Swiss-Prot: P22648]: FGFR
binding_
349      motif
351 <400> SEQUENCE: 16
353 Ala Thr Asn Lys Gly Gly Glu Val Lys Lys Asn Gly His Leu
354 1          5          10
357 <210> SEQ ID NO: 17
359 <211> LENGTH: 16
361 <212> TYPE: PRT
363 <213> ORGANISM: Artificial sequence
367 <220> FEATURE:
369 <223> OTHER INFORMATION: ADAM-19 precursor (EC 3.4.24.-) [Swiss-Prot: Q9H013/O35674]:
FGFR
370      binding motif
372 <400> SEQUENCE: 17
374 Lys Tyr Val Glu Leu Tyr Leu Val Ala Asp Tyr Leu Glu Phe Gln Lys
375 1          5          10          15
378 <210> SEQ ID NO: 18
380 <211> LENGTH: 15
382 <212> TYPE: PRT
384 <213> ORGANISM: Artificial sequence
388 <220> FEATURE:
390 <223> OTHER INFORMATION: ADAM-8 precursor (EC 3.4.24.-) [Swiss-Prot: P78325]: FGFR
binding
391      motif
393 <400> SEQUENCE: 18
395 Arg Tyr Val Glu Leu Tyr Val Val Val Asp Asn Ala Glu Phe Gln
396 1          5          10          15
399 <210> SEQ ID NO: 19
401 <211> LENGTH: 16
403 <212> TYPE: PRT
405 <213> ORGANISM: Artificial sequence
409 <220> FEATURE:
411 <223> OTHER INFORMATION: ADAM-12 precursor (EC 3.4.24.-) [Swiss-Prot: O43184; Q61824]:
FGFR
412      binding motif

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414 <400> SEQUENCE: 19



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Input Set : A:\CorrectedSEQ list P770PC00.ST25.txt

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416 Lys Tyr Val Glu Leu Val Ile Val Ala Asp Asn Arg Glu Phe Gln Arg
417 1          5          10          15
420 <210> SEQ ID NO: 20
422 <211> LENGTH: 16
424 <212> TYPE: PRT
426 <213> ORGANISM: Artificial sequence
430 <220> FEATURE:
432 <223> OTHER INFORMATION: Metalloproteinase-disintegrin domain containing protein
TECADAM [
433      AF163291] : FGFR binding motif
435 <400> SEQUENCE: 20
437 Lys Tyr Ile Glu Tyr Tyr Val Val Leu Asp Asn Gly Glu Phe Lys Lys
438 1          5          10          15
441 <210> SEQ ID NO: 21
443 <211> LENGTH: 14
445 <212> TYPE: PRT
447 <213> ORGANISM: Artificial sequence
451 <220> FEATURE:
453 <223> OTHER INFORMATION: ADAM-33 precursor (EC 3.4.24. -) [Swiss-Prot: Q9BZ11/Q923W9]
FGFR
454      binding motif
456 <400> SEQUENCE: 21
458 Arg Tyr Leu Glu Leu Tyr Ile Val Ala Asp His Thr Leu Phe
459 1          5          10
462 <210> SEQ ID NO: 22
464 <211> LENGTH: 15
466 <212> TYPE: PRT
468 <213> ORGANISM: Artificial sequence
472 <220> FEATURE:
474 <223> OTHER INFORMATION: ADAM-1A Fertilin alpha [Swiss-Prot: Q8R533]: FGFR binding
motif
476 <400> SEQUENCE: 22
478 Lys Tyr Val Glu Met Phe Val Val Val Asn His Gln Arg Phe Gln
479 1          5          10          15
482 <210> SEQ ID NO: 23
484 <211> LENGTH: 14
486 <212> TYPE: PRT
488 <213> ORGANISM: Artificial sequence
492 <220> FEATURE:
494 <223> OTHER INFORMATION: ADAM-9 [Swiss-Prot: Q13433; Q61072]: FGFR binding motif
496 <400> SEQUENCE: 23
498 Arg Tyr Val Glu Leu Phe Ile Val Val Asp Lys Glu Arg Tyr
499 1          5          10
502 <210> SEQ ID NO: 24
504 <211> LENGTH: 16
506 <212> TYPE: PRT
508 <213> ORGANISM: Artificial sequence
512 <220> FEATURE:
514 <223> OTHER INFORMATION: ADAM-7 precursor [Swiss-Prot: Q9H2U9]: FGFR binding motif
516 <400> SEQUENCE: 24
518 Lys Tyr Val Glu Leu Phe Ile Val Ala Asp Asp Thr Val Tyr Arg Arg
519 1          5          10          15

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**VERIFICATION SUMMARY**

DATE: 02/13/2006

PATENT APPLICATION: US/10/567,365

TIME: 12:42:57

Input Set : A:\CorrectedSEQ list P770PC00.ST25.txt

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L:17 M:270 C: Current Application Number differs, Replaced Current Application No

L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date